

**【CLAIMS】****【Claim 1】**

An isolated gene regulating fruit and seed development selected from a group consisting of a gene having a nucleotide sequence set forth in SEQ. ID. No 1 containing a nucleotide sequence encoding MADS-domain, a gene having a nucleotide sequence set forth in SEQ. ID. No 2 containing a nucleotide sequence encoding MADS-domain and a gene encoding an amino acid sequence having at least 85% homology within the region other than MADS-domain.

**【Claim 2】**

An expression vector comprising the gene according to Claim 1.

**【Claim 3】**

The expression vector according to Claim 2 wherein the expression vector is *pMdMADS14* into which a gene having the nucleotide sequence set forth in SEQ. ID. No 1 is inserted in forward direction (Accession No: KCTC 10588BP).

**【Claim 4】**

The expression vector according to Claim 2 wherein the expression vector is *pMdMADS16* into which a gene having the nucleotide sequence set forth in SEQ. ID. No 2 is inserted in forward direction (Accession No: KCTC 10589BP).

**【Claim 5】**

A transgenic plant cell containing the gene according to Claim 1.

**【Claim 6】**

5        A transgenic plant whose fruit and seed development is regulated, and that is prepared by regeneration of the transgenic plant cells according to Claim 5 by tissue culture technique.

**【Claim 7】**

10        The transgenic plant according to Claim 6 wherein the plant is selected from a group consisting of food crops such as rice, wheat, barley, corns, soybean, potato, red bean, oat, sorghum; vegetables such as Chinese cabbage, radish, red pepper, strawberry, tomato, watermelon,  
15        cucumber, cabbage, melon, pumpkin, spring onion, onion, carrot; industrial crops such as ginseng, *Acanthopanax senticosus*, tobacco, cotton, sesame, sugar cane, sugar beet, *Perilla japonica*, peanut, rape; fruits such as apple, pear, orange, jujube, peach, kiwifruit, grapes, tangerine,  
20        persimmon, plum, apricot, bananas; floricultural crops such as rose, gladiolus, gerbera, carnation, chrysanthemum, lily, tulip; forage crops such as ryegrass, red clover, orchard grass, alfalfa, tall fescue, perennial ryegrass; fiber crops such as cotton plant; and landscape plants  
25        such as flowers and shrubs.

**【Claim 8】**

An offspring or a clone of a transgenic plant according to Claim 6.

**5      【Claim 9】**

A Fruit, seed, ear, tuber, tuberos root, column, callus or a protoplast of a transgenic plant according to Claim 6.

**10      【Claim 10】**

The transgenic plant according to Claim 6 wherein the plant shows one of the following phenotypes:

15      a phenotype in which sepal is transformed into fruit flesh and parthenocarpic fruit is formed; a phenotype in which seed development is promoted and ripening is delayed; and a phenotype in which fruit and seed development is inhibited.

**20      【Claim 11】**

A method of preparing a transgenic plant whose fruit and seed development was regulated, comprising the steps of:

1) Constructing an expression vector comprising the gene according to Claim 1;

2) Transferring the vector constructed in Step 1) into Agrobacterium;

3) Co-culturing the transformed Agrobacterium of step 2) with plant tissue; and

4) Regenerating the transformed tissue into a mature transgenic plant.

5     **【Claim 12】**

A Composition for fruit and seed development in a plant comprising the gene according to Claim 1 or the expression vector according to any one of Claims 2-4 as an effective ingredient.

10    **【Claim 13】**

A Composition for regulating the synthesis of active gibberellin containing the gene according to Claim 1 or the expression vector according to any one of Claims 2-4 as an effective ingredient.

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